

Supplementary Data Set. Sequences that were retrieved with significant e-values using BAE61530 as the query for a BLAST search against the non-redundant protein sequence database of the NCBI

GenBank accession	protein name in GenBank
BAE61530.1	unnamed protein product [Aspergillus oryzae RIB40]
XP_002382772.1	conserved hypothetical protein [Aspergillus flavus NRRL3357]
EIT79819.1	hypothetical protein Ao3042_03704 [Aspergillus oryzae 3.042]
EHA28482.1	hypothetical protein ASPNIDRAFT_202949 [Aspergillus niger ATCC 1015]
XP_001393828.1	hypothetical protein ANI_1_634084 [Aspergillus niger CBS 513.88] >emb CAK40369.1 unnamed protein product [Aspergillus niger]
GAA88290.1	endoglucanase [Aspergillus kawachii IFO 4308]
XP_001276229.1	conserved hypothetical protein [Aspergillus clavatus NRRL 1] >gb EAW14803.1 conserved hypothetical protein [Aspergillus clavatus NRRL 1]
XP_001266231.1	hypothetical protein NFIA_039100 [Neosartorya fischeri NRRL 181] >gb EAW24334.1 conserved hypothetical protein [Neosartorya fischeri NRRL 181]
XP_748042.1	conserved hypothetical protein [Aspergillus fumigatus Af293] >gb EAL86004.1 conserved hypothetical protein [Aspergillus fumigatus Af293]
EDP51150.1	conserved hypothetical protein [Aspergillus fumigatus A1163]
XP_001217899.1	predicted protein [Aspergillus terreus NIH2624] >gb EAU30414.1 predicted protein [Aspergillus terreus NIH2624]
XP_002568660.1	Pc21g16530 [Penicillium chrysogenum Wisconsin 54-1255] >emb CAP96550.1 Pc21g16530 [Penicillium chrysogenum Wisconsin 54-1255]
EKV10928.1	hypothetical protein PDIG_54160 [Penicillium digitatum PHI26] >gb EKV13250.1 hypothetical protein PDIP_49380 [Penicillium digitatum Pd1]
XP_002481851.1	conserved hypothetical protein [Talaromyces stipitatus ATCC 10500] >gb EED17859.1 conserved hypothetical protein [Talaromyces stipitatus ATCC 10500]
XP_002147680.1	conserved hypothetical protein [Talaromyces marneffeii ATCC 18224] >gb EEA24169.1 conserved hypothetical protein [Talaromyces marneffeii ATCC 18224]
EPS34477.1	chitin binding domain-containing protein [Penicillium oxalicum 114-2]
XP_001242606.1	hypothetical protein CIMG_06502 [Coccidioides immitis RS] >gb EAS31305.2 hypothetical protein CIMG_06502 [Coccidioides immitis RS]
XP_003069792.1	hypothetical protein CPC735_029830 [Coccidioides posadasii C735 delta SOWgp]
XP_002566969.1	Pc17g00010 [Penicillium chrysogenum Wisconsin 54-1255] >emb CAP79288.1 Pc17g00010, partial [Penicillium chrysogenum Wisconsin 54-1255]
XP_002556886.1	Pc09g00200 [Penicillium chrysogenum Wisconsin 54-1255] >emb CAP79267.1 Pc09g00200 [Penicillium chrysogenum Wisconsin 54-1255]
EHY55011.1	DNA-directed RNA polymerase [Exophiala dermatitidis NIH/UT8656]
XP_002544405.1	predicted protein [Uncinocarpus reesii 1704] >gb EEP79076.1 predicted protein [Uncinocarpus reesii 1704]
XP_002566537.1	Pc22g26710 [Penicillium chrysogenum Wisconsin 54-1255] >emb CAP99959.1 Pc22g26710 [Penicillium chrysogenum Wisconsin 54-1255]
XP_002566575.1	Pc23g00050 [Penicillium chrysogenum Wisconsin 54-1255] >emb CAP79499.1 Pc23g00050 [Penicillium chrysogenum Wisconsin 54-1255]
XP_002563773.1	Pc20g12880 [Penicillium chrysogenum Wisconsin 54-1255] >emb CAP86617.1 Pc20g12880 [Penicillium chrysogenum Wisconsin 54-1255]
XP_003174002.1	hypothetical protein MGVY_04176 [Arthroderma gypseum CBS 118893] >gb EFR01172.1 hypothetical protein MGVY_04176 [Arthroderma gypseum CBS 118893]
EKG18934.1	hypothetical protein MPH_03818 [Macrophomina phaseolina MS6]
XP_002848955.1	conserved hypothetical protein [Arthroderma otae CBS 113480] >gb EEQ29070.1 conserved hypothetical protein [Arthroderma otae CBS 113480]
XP_003015133.1	conserved hypothetical protein [Arthroderma benhamiae CBS 112371] >gb EFE34493.1 conserved hypothetical protein [Arthroderma benhamiae CBS 112371]
XP_003019976.1	conserved hypothetical protein [Trichophyton verrucosum HKI 0517] >gb EFE39352.1 conserved hypothetical protein [Trichophyton verrucosum HKI 0517]
XP_001597818.1	hypothetical protein SS1G_02014 [Sclerotinia sclerotiorum 1980] >gb EDN97086.1 hypothetical protein SS1G_02014 [Sclerotinia sclerotiorum 1980 UF-70]
EGD92942.1	hypothetical protein TESHG_00502 [Trichophyton tonsurans CBS 112818]
EGE04098.1	extracellular protein [Trichophyton equinum CBS 127.97]
XP_003235451.1	hypothetical protein TERG_04504 [Trichophyton rubrum CBS 118892] >gb EGD88256.1 hypothetical protein TERG_04504 [Trichophyton rubrum CBS 118892]
EME82432.1	hypothetical protein MYCFIDRAFT_59548 [Pseudocercospora fijiensis CIRAD86]
XP_002789213.1	conserved hypothetical protein [Paracoccidioides sp. 'lutzii' Pb01] >gb EEH40026.1 conserved hypothetical protein [Paracoccidioides sp. 'lutzii' Pb01]
ELR02541.1	hypothetical protein GMDG_01066 [Pseudogymnoascus destructans 20631-21]
XP_003050482.1	hypothetical protein NECHADRAFT_96270 [Nectria haematococca mpVI 77-13-4] >gb EEU44769.1 hypothetical protein NECHADRAFT_96270 [Nectria haematococca mpVI 77-13-4]
EEH05325.1	conserved hypothetical protein [Ajellomyces capsulatus G186AR]
XP_001543425.1	predicted protein [Ajellomyces capsulatus NAM1] >gb EDN02607.1 predicted protein [Ajellomyces capsulatus NAM1]
EGC46978.1	conserved hypothetical protein [Ajellomyces capsulatus H88]
EEH49643.1	conserved hypothetical protein [Paracoccidioides brasiliensis Pb18]
EME42943.1	hypothetical protein DOTSEDRAFT_72390 [Dothistroma septosporum NZE10]
EEH23200.1	conserved hypothetical protein [Paracoccidioides brasiliensis Pb03]
XP_681444.1	hypothetical protein AN8175.2 [Aspergillus nidulans FGSC A4]
CCD52974.1	hypothetical protein BofuT4P90000026001 [Botryotinia fuckeliana T4]
XP_001547004.1	hypothetical protein BC1G_14469 [Botryotinia fuckeliana B05.10]
EMR83340.1	putative chitin- domain 3 protein [Botryotinia fuckeliana BcdW1]
CAQ16288.1	hypothetical protein [Colletotrichum graminicola]
EHK20062.1	hypothetical protein TRIVIDRAFT_171588, partial [Trichoderma virens Gv29-8]
EFQ29865.1	hypothetical protein GLRG_05009 [Colletotrichum graminicola M1.001]

EMF11929.1 hypothetical protein SEPMUDRAFT_142137 [Sphaerulina musiva SO2202]
 EON63048.1 hypothetical protein W97_02275 [Coniosporium apollinis CBS 100218]
 XP_003850030.1 hypothetical protein MYCGRDRAFT_46866 [Zymoseptoria tritici IPO323] >gb|EGP85006.1| hypothetical protein MYCGRDRAFT_46866 [Zymoseptoria tritici IPO323]
 ELA36568.1 hypothetical protein CGGC5_3835 [Colletotrichum gloeosporioides Nara gc5]
 EGE82146.1 hypothetical protein BDDG_05089 [Ajellomyces dermatitidis ATCC 18188]
 EQL33487.1 hypothetical protein BDFG_04625 [Ajellomyces dermatitidis ATCC 26199]
 XP_002628516.1 conserved hypothetical protein [Ajellomyces dermatitidis SLH14081] >gb|EEQ73194.1| conserved hypothetical protein [Ajellomyces dermatitidis SLH14081]
 EMD86748.1 hypothetical protein COCHEDRAFT_1115227 [Bipolaris maydis C5] >gb|ENI03140.1| hypothetical protein COCC4DRAFT_143749 [Bipolaris maydis ATCC 48331]
 CCF45557.1 endoglucanase [Colletotrichum higginsianum]
 ENH82730.1 glycoside hydrolase [Colletotrichum orbiculare MAFF 240422]
 EHK42112.1 hypothetical protein TRIATDRAFT_302363 [Trichoderma atroviride IMI 206040]
 XP_003307044.1 hypothetical protein PTT_20365 [Pyrenophora teres f. teres 0-1] >gb|EFQ84871.1| hypothetical protein PTT_20365 [Pyrenophora teres f. teres 0-1]
 XP_001941363.1 conserved hypothetical protein [Pyrenophora tritici-repentis Pt-1C-BFP] >gb|EDU44082.1| conserved hypothetical protein [Pyrenophora tritici-repentis Pt-1C-BFP]
 EON61299.1 hypothetical protein W97_00512 [Coniosporium apollinis CBS 100218]
 EGR49412.1 glycoside hydrolase [Trichoderma reesei QM6a]
 CCT67099.1 related to spore coat protein SP96 precursor [Fusarium fujikuroi IMI 58289]
 ERF75887.1 hypothetical protein EPUS_01253 [Endocarpon pusillum Z07020]
 EJP64324.1 Chitin-binding, domain 3 [Beauveria bassiana ARSEF 2860]
 EPE05022.1 glycoside hydrolase [Ophiostoma piceae UAMH 11346]
 EFZ01910.1 hypothetical protein MAA_03139 [Metarhizium anisopliae ARSEF 23]
 CCE27776.1 related to spore coat protein SP96 precursor [Claviceps purpurea 20.1]
 EPE24813.1 hypothetical protein GLAREA_11394 [Glarea lozoyensis ATCC 20868]
 EOA84296.1 hypothetical protein SETTUDRAFT_164498 [Setosphaeria turcica Et28A]
 EFX02550.1 hypothetical protein CMQ_5911 [Grosmanina clavigera kw1407]
 EME49105.1 hypothetical protein DOTSEDRAFT_49429 [Dothiostroma septosporum NZE10]
 XP_003006162.1 conserved hypothetical protein [Verticillium alfalfae VaMs.102] >gb|EEY18006.1| conserved hypothetical protein [Verticillium alfalfae VaMs.102]
 EGY17657.1 hypothetical protein VDAG_01339 [Verticillium dahliae VdLs.17]
 EMT68301.1 hypothetical protein FOC4_g10012714 [Fusarium oxysporum f. sp. cubense race 4]
 EMD67171.1 hypothetical protein COCSADRAFT_285519 [Bipolaris sorokiniana ND90Pr]
 EKD14716.1 hypothetical protein MBM_06927 [Marssonina brunnea f. sp. 'multigermtubi' MB_m1]
 ENH68621.1 hypothetical protein FOC1_g10014406 [Fusarium oxysporum f. sp. cubense race 1]
 EGU85465.1 hypothetical protein FOXB_03949 [Fusarium oxysporum Fo5176]
 EMT68300.1 hypothetical protein FOC4_g10012713 [Fusarium oxysporum f. sp. cubense race 4]
 EGX94965.1 hypothetical protein CCM_03237 [Cordyceps militaris CM01]
 EKJ75600.1 hypothetical protein FPSE_04243 [Fusarium pseudograminearum CS3096]
 XP_386263.1 hypothetical protein FG06087.1 [Fusarium graminearum PH-1]
 EMD61301.1 hypothetical protein COCSADRAFT_60631, partial [Bipolaris sorokiniana ND90Pr]
 EQL04114.1 glycoside hydrolase [Ophiocordyceps sinensis CO18]
 EMD86119.1 hypothetical protein COCHEDRAFT_1023915 [Bipolaris maydis C5]
 EON97679.1 putative chitin- domain 3 protein [Togninia minima UCRPA7]
 XP_001228963.1 hypothetical protein CHGG_02447 [Chaetomium globosum CBS 148.51]
 XP_960937.1 hypothetical protein NCU01380 [Neurospora crassa OR74A]
 ENI08410.1 hypothetical protein COCC4DRAFT_110653, partial [Bipolaris maydis ATCC 48331]
 EMD91831.1 hypothetical protein COCHEDRAFT_1214230 [Bipolaris maydis C5]
 EFY91792.1 hypothetical protein MAC_02077 [Metarhizium acridum CQM a 102]
 EFZ04469.1 hypothetical protein MAA_01543 [Metarhizium anisopliae ARSEF 23]
 EPE10862.1 dna-directed rna polymerase [Ophiostoma piceae UAMH 11346]
 EPS39968.1 hypothetical protein H072_6402 [Dactylellina haptotyla CBS 200.50]
 XP_001931006.1 conserved hypothetical protein [Pyrenophora tritici-repentis Pt-1C-BFP]
 EOA82325.1 hypothetical protein SETTUDRAFT_73750, partial [Setosphaeria turcica Et28A]
 EEQ89326.1 conserved hypothetical protein [Ajellomyces dermatitidis ER-3]
 CCU83020.1 hypothetical protein BGHHD14_bgh02286 [Blumeria graminis f. sp. hordei DH14]
 XP_003305261.1 hypothetical protein PTT_18064 [Pyrenophora teres f. teres 0-1]

EPQ64850.1 hypothetical protein BGT96224_2589 [Blumeria graminis f. sp. tritici 96224]
EMR69119.1 putative dna-directed rna polymerase protein [Eutypa lata UCREL1]
EGO60701.1 hypothetical protein NEUTE1DRAFT_127516 [Neurospora tetrasperma FGSC 2508]
ELR06576.1 hypothetical protein GMDG_08049 [Pseudogymnoascus destructans 20631-21]
XP_003348222.1 hypothetical protein SMAC_04067 [Sordaria macrospora k-hell] >emb|CCC11364.1| unnamed protein product [Sordaria macrospora k-hell]
XP_001911666.1 hypothetical protein [Podospora anserina S mat+] >emb|CAP73493.1| unnamed protein product [Podospora anserina S mat+]
XP_003836245.1 predicted protein [Leptosphaeria maculans JN3] >emb|CBX92880.1| predicted protein [Leptosphaeria maculans JN3]
XP_002562514.1 Pc19g00200 [Penicillium chrysogenum Wisconsin 54-1255] >emb|CAP79436.1| Pc19g00200 [Penicillium chrysogenum Wisconsin 54-1255]
EMF17657.1 hypothetical protein SEPMUDRAFT_35015, partial [Sphaerulina musiva SO2202]
EJT69668.1 hypothetical protein GGTG_12552 [Gaeumannomyces graminis var. tritici R3-111a-1]
EMD65510.1 hypothetical protein COCSADRAFT_189304 [Bipolaris sorokiniana ND90Pr]
EFW99073.1 hypothetical protein CMQ_5494 [Grosmanina clavigera kw1407]
CCF32152.1 endoglucanase [Colletotrichum higginsianum]
EMR68901.1 putative dna-directed rna polymerase protein [Eutypa lata UCREL1]
EOD43831.1 putative dna-directed rna polymerase protein [Neofusicoccum parvum UCRNP2]
EMR69820.1 putative chitin- domain 3 protein [Eutypa lata UCREL1]
XP_001937368.1 conserved hypothetical protein [Pyrenophora tritici-repentis Pt-1C-BFP]
EFQ36410.1 endoglucanase [Colletotrichum graminicola M1.001]
EOD46824.1 hypothetical protein UCRNP2_6442 [Neofusicoccum parvum UCRNP2]
XP_003303971.1 hypothetical protein PTT_16375 [Pyrenophora teres f. teres 0-1] >gb|EFQ87921.1| hypothetical protein PTT_16375 [Pyrenophora teres f. teres 0-1]
EMD01144.1 hypothetical protein BAUCODRAFT_52546, partial [Baudoinia compniacensis UAMH 10762]
XP_003856632.1 hypothetical protein MYCGRDRAFT_32157, partial [Zymoseptoria tritici IPO323]
ENH83182.1 glycoside hydrolase [Colletotrichum orbiculare MAFF 240422]
EMD66052.1 hypothetical protein COCSADRAFT_355596 [Bipolaris sorokiniana ND90Pr]
XP_003349120.1 hypothetical protein SMAC_06957 [Sordaria macrospora k-hell] >emb|CCC12549.1| unnamed protein product [Sordaria macrospora k-hell]
XP_003712213.1 hypothetical protein MGG_09527 [Magnaporthe oryzae 70-15] >gb|EHA52406.1| hypothetical protein MGG_09527 [Magnaporthe oryzae 70-15]
ELQ38216.1 hypothetical protein OOU_Y34scaffold00548g32 [Magnaporthe oryzae Y34]
ELQ65882.1 hypothetical protein OOW_P131scaffold00450g11 [Magnaporthe oryzae P131]
EGX53029.1 hypothetical protein AOL_s00007g365 [Arthrobotrys oligospora ATCC 24927]
EGO54744.1 hypothetical protein NEUTE1DRAFT_141028 [Neurospora tetrasperma FGSC 2508]
EMR67112.1 putative chitin- domain 3 protein [Eutypa lata UCREL1]
XP_958127.1 hypothetical protein NCU05932 [Neurospora crassa OR74A] >gb|EAA28891.1| predicted protein [Neurospora crassa OR74A]
XP_001791303.1 hypothetical protein SNOG_00622 [Phaeosphaeria nodorum SN15] >gb|EAT92117.1| hypothetical protein SNOG_00622 [Phaeosphaeria nodorum SN15]
EGZ67778.1 hypothetical protein NEUTE2DRAFT_28764, partial [Neurospora tetrasperma FGSC 2509]
XP_001801474.1 hypothetical protein SNOG_11232 [Phaeosphaeria nodorum SN15] >gb|EAT81731.2| hypothetical protein SNOG_11232 [Phaeosphaeria nodorum SN15]
XP_003658112.1 hypothetical protein THITE_2124610 [Thielavia terrestris NRRL 8126] >gb|AEO71776.1| hypothetical protein THITE_2124610 [Thielavia terrestris NRRL 8126]
EOA81334.1 hypothetical protein SETTUDRAFT_165829 [Setosphaeria turcica Et28A]
EQB46896.1 hypothetical protein CGLO_14025 [Colletotrichum gloeosporioides Cg-14]
EEQ83434.1 endoglucanase [Ajellomyces dermatitidis ER-3] >gb|EQL35978.1| DNA-directed RNA polymerase [Ajellomyces dermatitidis ATCC 26199]
XP_001214248.1 conserved hypothetical protein [Aspergillus terreus NIH2624] >gb|EAU34139.1| conserved hypothetical protein [Aspergillus terreus NIH2624]
EQL01822.1 hypothetical protein OCS_02460 [Ophiocordyceps sinensis CO18]
EJT82590.1 hypothetical protein GGTG_02563 [Gaeumannomyces graminis var. tritici R3-111a-1]
XP_003661558.1 hypothetical protein MYCTH_2301079 [Myceliophthora thermophila ATCC 42464] >gb|AEO56313.1| hypothetical protein MYCTH_2301079 [Myceliophthora thermophila ATCC 42464]
XP_002621567.1 endoglucanase [Ajellomyces dermatitidis SLH14081] >gb|EEQ73976.1| endoglucanase [Ajellomyces dermatitidis SLH14081]
XP_001542256.1 predicted protein [Ajellomyces capsulatus NAM1] >gb|EDN05824.1| predicted protein [Ajellomyces capsulatus NAM1]
EFZ02283.1 extracellular protein, putative [Metarhizium anisopliae ARSEF 23]
XP_001909751.1 hypothetical protein [Podospora anserina S mat+] >emb|CAP70884.1| unnamed protein product [Podospora anserina S mat+]
ELA28398.1 endoglucanase, partial [Colletotrichum gloeosporioides Nara gc5]
EFY88319.1 RNA polymerase Rpb1 repeat domain protein [Metarhizium acridum CQMa 102]
ELA24573.1 hypothetical protein CGGC5_2157 [Colletotrichum gloeosporioides Nara gc5]
XP_003021860.1 hypothetical protein TRV_04037 [Trichophyton verrucosum HKI 0517] >gb|EFE41242.1| hypothetical protein TRV_04037 [Trichophyton verrucosum HKI 0517]
CCT75452.1 related to spore coat protein SP96 precursor [Fusarium fujikuroi IMI 58289]
XP_003660925.1 hypothetical protein MYCTH_2299757 [Myceliophthora thermophila ATCC 42464] >gb|AEO55680.1| hypothetical protein MYCTH_2299757 [Myceliophthora thermophila ATCC 42464]

EGC44326.1 endoglucanase [Ajellomyces capsulatus H88]

EEH09854.1 conserved hypothetical protein [Ajellomyces capsulatus G186AR]

XP_002835569.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ79726.1| unnamed protein product [Tuber melanosporum]

XP_002487567.1 extracellular protein, putative [Talaromyces stipitatus ATCC 10500] >gb|EED13456.1| extracellular protein, putative [Talaromyces stipitatus ATCC 10500]

EEQ89863.1 endoglucanase [Ajellomyces dermatitidis ER-3]

EGE80900.1 endoglucanase [Ajellomyces dermatitidis ATCC 18188]

XP_002621360.1 endoglucanase [Ajellomyces dermatitidis SLH14081] >gb|EEQ74177.1| endoglucanase [Ajellomyces dermatitidis SLH14081]

EER38195.1 endoglucanase [Ajellomyces capsulatus H143]

EGE09367.1 endoglucanase [Trichophyton equinum CBS 127.97]

ERF68296.1 hypothetical protein EPUS_02752 [Endocarpon pusillum Z07020]

XP_002153103.1 extracellular protein, putative [Talaromyces marneffeii ATCC 18224] >gb|EEA18718.1| extracellular protein, putative [Talaromyces marneffeii ATCC 18224]

XP_658206.1 hypothetical protein AN0602.2 [Aspergillus nidulans FGSC A4]

XP_002540653.1 predicted protein [Uncinocarpus reesii 1704] >gb|EEP75320.1| predicted protein [Uncinocarpus reesii 1704]

EDP49665.1 extracellular protein, putative [Aspergillus fumigatus A1163]

XP_001258215.1 RNA polymerase Rpb1 C-terminal repeat domain protein [Neosartorya fischeri NRRL 181] >gb|EAW16318.1| RNA polymerase Rpb1 C-terminal repeat domain protein [Neosartorya fischeri NRRL 181]

XP_748715.1 endoglucanase [Aspergillus fumigatus Af293] >gb|EAL86677.1| endoglucanase, putative [Aspergillus fumigatus Af293] >gb|EDP53229.1| endoglucanase, putative [Aspergillus fumigatus A1163]

EGE03336.1 endoglucanase [Trichophyton equinum CBS 127.97]

EMF11619.1 hypothetical protein SEPMUDRAFT_10541, partial [Sphaerulina musiva SO2202]

EMD89928.1 hypothetical protein COCHEDRAFT_1178048 [Bipolaris maydis CS] >gb|ENI09860.1| hypothetical protein COCC4DRAFT_185311 [Bipolaris maydis ATCC 48331]

EME82188.1 hypothetical protein MYCFIDRAFT_215607 [Pseudocercospora fijiensis CIRAD86]

XP_001259157.1 endoglucanase, putative [Neosartorya fischeri NRRL 181] >gb|EAW17260.1| endoglucanase, putative [Neosartorya fischeri NRRL 181]

EQL37881.1 DNA-directed RNA polymerase [Ajellomyces dermatitidis ATCC 26199]

EGR50937.1 predicted protein [Trichoderma reesei QM6a]

XP_002380776.1 endoglucanase, putative [Aspergillus flavus NRRL3357] >gb|EED48875.1| endoglucanase, putative [Aspergillus flavus NRRL3357] >gb|EIT81221.1| endoglucanase, putative [Aspergillus oryzae 3.042]

XP_750980.1 extracellular protein [Aspergillus fumigatus Af293] >gb|EAL88942.1| extracellular protein, putative [Aspergillus fumigatus Af293]

XP_001268194.1 conserved hypothetical protein [Aspergillus clavatus NRRL 1] >gb|EAW06768.1| conserved hypothetical protein [Aspergillus clavatus NRRL 1]

XP_003013767.1 hypothetical protein ARB_07879 [Arthroderma benhamiae CBS 112371] >gb|EFE33127.1| hypothetical protein ARB_07879 [Arthroderma benhamiae CBS 112371]

XP_003837919.1 hypothetical protein LEMA_P119660.1 [Leptosphaeria maculans JN3] >emb|CBX94475.1| hypothetical protein LEMA_P119660.1 [Leptosphaeria maculans JN3]

EKJ75109.1 hypothetical protein FPSE_04729 [Fusarium pseudograminearum CS3096]

EGS21088.1 hypothetical protein CHTT_0029290 [Chaetomium thermophilum var. thermophilum DSM 1495]

EKD20942.1 hypothetical protein MBM_00055 [Marssonina brunnea f. sp. 'multigermtubi' MB_m1]

EFY86242.1 extracellular protein, putative [Metarhizium acridium CQMa 102]

XP_003024865.1 hypothetical protein TRV_00940 [Trichophyton verrucosum HKI 0517] >gb|EFE44254.1| hypothetical protein TRV_00940 [Trichophyton verrucosum HKI 0517]

EME43051.1 hypothetical protein DOTSEDRAFT_53974 [Dothistroma septosporum NZE10]

CCT62801.1 related to spore coat protein SP96 precursor [Fusarium fujikuroi IMI 58289]

E0001866.1 putative spore coat protein sp96 precursor-like protein [Togninia minima UCRPA7]

EGD93178.1 endoglucanase [Trichophyton tonsurans CBS 112818]

XP_001823731.1 endoglucanase [Aspergillus oryzae RIB40] >dbj|BAE62598.1| unnamed protein product [Aspergillus oryzae RIB40]

XP_003231518.1 endoglucanase [Trichophyton rubrum CBS 118892] >gb|EGD91599.1| endoglucanase [Trichophyton rubrum CBS 118892]

EGU83897.1 hypothetical protein FOXB_05611 [Fusarium oxysporum Fo5176]

EMT61439.1 hypothetical protein FOC4_g10014498 [Fusarium oxysporum f. sp. cubense race 4]

ENH61451.1 hypothetical protein FOC1_g10016267 [Fusarium oxysporum f. sp. cubense race 1]

EKG19379.1 hypothetical protein MPH_03242 [Macrophomina phaseolina MS6]

XP_003235677.1 endoglucanase [Trichophyton rubrum CBS 118892] >gb|EGD86472.1| endoglucanase [Trichophyton rubrum CBS 118892]

EGD98498.1 endoglucanase [Trichophyton tonsurans CBS 112818]

EMT69478.1 hypothetical protein FOC4_g10003203 [Fusarium oxysporum f. sp. cubense race 4]

ENH71945.1 hypothetical protein FOC1_g10002138 [Fusarium oxysporum f. sp. cubense race 1]

EJP65660.1 extracellular protein [Beauveria bassiana ARSEF 2860]

XP_380818.1 hypothetical protein FG00642.1 [Fusarium graminearum PH-1]

XP_001396774.1 extracellular protein [Aspergillus niger CBS 513.88] >emb|CAK42338.1| unnamed protein product [Aspergillus niger]

XP_001393259.2 extracellular protein [Aspergillus niger CBS 513.88]

EQB58330.1 endoglucanase [Colletotrichum gloeosporioides Cg-14]

XP_003174226.1 hypothetical protein MGYG_04403 [Arthroderma gypseum CBS 118893] >gb|EFR01396.1| hypothetical protein MGYG_04403 [Arthroderma gypseum CBS 118893]

XP_003856402.1 hypothetical protein MYCGRDRAFT_33493, partial [Zyloseptoria tritici IPO323] >gb|EGP91378.1| hypothetical protein MYCGRDRAFT_33493 [Zyloseptoria tritici IPO323]
 EGE05564.1 extracellular protein [Trichophyton equinum CBS 127.97]
 XP_002792749.1 conserved hypothetical protein [Paracoccidioides sp. 'lutzii' Pb01] >gb|EEH34429.1| conserved hypothetical protein [Paracoccidioides sp. 'lutzii' Pb01]
 XP_003650198.1 hypothetical protein THITE_2023982, partial [Thielavia terrestris NRRL 8126] >gb|AEO63862.1| hypothetical protein THITE_2023982, partial [Thielavia terrestris NRRL 8126]
 GAA90553.1 hypothetical protein AKAW_08667 [Aspergillus kawachii IFO 4308]
 EPE26388.1 endoglucanase [Glarea lozoyensis ATCC 20868]
 EHA24588.1 hypothetical protein ASPNIDRAFT_48700 [Aspergillus niger ATCC 1015]
 EGR52032.1 predicted protein [Trichoderma reesei QM6a]
 EKG12761.1 hypothetical protein MPH_10105 [Macrophomina phaseolina MS6]
 GAA93289.1 RNA polymerase Rpb1 C-terminal repeat domain protein [Aspergillus kawachii IFO 4308]
 XP_003053715.1 hypothetical protein NECHADRAFT_103191 [Nectria haematococca mpVI 77-13-4] >gb|EEU48002.1| hypothetical protein NECHADRAFT_103191 [Nectria haematococca mpVI 77-13-4]
 Efy94360.1 endoglucanase, putative [Metarhizium anisopliae ARSEF 23]
 XP_002482903.1 endoglucanase, putative [Talaromyces stipitatus ATCC 10500] >gb|EED18911.1| endoglucanase, putative [Talaromyces stipitatus ATCC 10500]
 EMC93960.1 hypothetical protein BAUCODRAFT_216599 [Baudoinia compniacensis UAMH 10762]
 EGX91035.1 extracellular protein, putative [Cordyceps militaris CM01]
 XP_003033631.1 hypothetical protein SCHCODRAFT_256354 [Schizophyllum commune H4-8] >gb|EFI98728.1| hypothetical protein SCHCODRAFT_256354 [Schizophyllum commune H4-8]
 EKJ73219.1 hypothetical protein FPSE_06643 [Fusarium pseudograminearum CS3096]
 EGU87901.1 hypothetical protein FOXB_01587 [Fusarium oxysporum Fo5176]
 EQL00501.1 hypothetical protein OCS_03783 [Ophiocordyceps sinensis CO18]
 XP_003169269.1 hypothetical protein MGYG_08815 [Arthroderma gypseum CBS 118893] >gb|EFQ96892.1| hypothetical protein MGYG_08815 [Arthroderma gypseum CBS 118893]
 XP_003032165.1 hypothetical protein SCHCODRAFT_85194 [Schizophyllum commune H4-8] >gb|EFI97262.1| hypothetical protein SCHCODRAFT_85194 [Schizophyllum commune H4-8]
 EGU76897.1 hypothetical protein FOXB_12585 [Fusarium oxysporum Fo5176]
 CCE28971.1 related to spore coat protein SP96 precursor [Claviceps purpurea 20.1]
 XP_003045352.1 hypothetical protein NECHADRAFT_44080 [Nectria haematococca mpVI 77-13-4] >gb|EEU39639.1| hypothetical protein NECHADRAFT_44080 [Nectria haematococca mpVI 77-13-4]
 EMT70525.1 hypothetical protein FOC4_g10008988 [Fusarium oxysporum f. sp. cubense race 4]
 CAK45829.1 unnamed protein product [Aspergillus niger]
 XP_002566688.1 Pc24g00280 [Penicillium chrysogenum Wisconsin 54-1255] >emb|CAP86936.1| Pc24g00280 [Penicillium chrysogenum Wisconsin 54-1255]
 XP_384016.1 hypothetical protein FG03840.1 [Fusarium graminearum PH-1]
 XP_001821065.1 extracellular protein [Aspergillus oryzae RIB40]
 XP_002376809.1 extracellular protein, putative [Aspergillus flavus NRRL3357]
 EIT74896.1 extracellular protein [Aspergillus oryzae 3.042]
 XP_003712585.1 hypothetical protein MGG_05070 [Magnaporthe oryzae 70-15] >gb|EHA52778.1| hypothetical protein MGG_05070 [Magnaporthe oryzae 70-15]
 EGS23085.1 hypothetical protein CHTT_0015740 [Chaetomium thermophilum var. thermophilum DSM 1495]
 XP_963492.1 hypothetical protein NCU05404 [Neurospora crassa OR74A] >emb|CAB97468.1| related to spore coat protein SP96 precursor [Neurospora crassa]
 CCE32966.1 related to spore coat protein SP96 precursor [Claviceps purpurea 20.1]
 XP_003047765.1 hypothetical protein NECHADRAFT_105522 [Nectria haematococca mpVI 77-13-4] >gb|EEU42052.1| hypothetical protein NECHADRAFT_105522 [Nectria haematococca mpVI 77-13-4]
 XP_002842272.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ86463.1| unnamed protein product [Tuber melanosporum]
 EGO56821.1 hypothetical protein NEUTE1DRAFT_117496 [Neurospora tetrasperma FGSC 2508] >gb|EGZ70289.1| hypothetical protein NEUTE2DRAFT_144984 [Neurospora tetrasperma FGSC 2509]
 EKG20672.1 Chitin-binding domain 3 [Macrophomina phaseolina MS6]
 CCT69101.1 related to spore coat protein SP96 precursor [Fusarium fujikuroi IMI 58289]
 ELQ43122.1 hypothetical protein OOU_Y34scaffold00174g87 [Magnaporthe oryzae Y34] >gb|ELQ68119.1| hypothetical protein OOW_P131scaffold00266g5 [Magnaporthe oryzae P131]
 XP_001273546.1 endoglucanase, putative [Aspergillus clavatus NRRL 1] >gb|EAW12120.1| endoglucanase, putative [Aspergillus clavatus NRRL 1]
 ENH74483.1 hypothetical protein FOC1_g10009901 [Fusarium oxysporum f. sp. cubense race 1]
 XP_003174561.1 hypothetical protein MGYG_02092 [Arthroderma gypseum CBS 118893] >gb|EFQ99078.1| hypothetical protein MGYG_02092 [Arthroderma gypseum CBS 118893]
 EGE03508.1 endoglucanase [Trichophyton equinum CBS 127.97]
 XP_001228385.1 hypothetical protein CHGG_10458 [Chaetomium globosum CBS 148.51] >gb|EAQ84054.1| hypothetical protein CHGG_10458 [Chaetomium globosum CBS 148.51]
 CCE28972.1 related to spore coat protein SP96 precursor [Claviceps purpurea 20.1]
 EAS31685.2 endoglucanase [Coccidioides immitis RS]
 EKJ77892.1 hypothetical protein FPSE_01985 [Fusarium pseudograminearum CS3096]
 XP_003070051.1 hypothetical protein CPC735_032420 [Coccidioides posadasii C735 delta SOWgp] >gb|EER27906.1| hypothetical protein CPC735_032420 [Coccidioides posadasii C735 delta SOWgp]
 EHK43388.1 hypothetical protein TRIATDRAFT_18328, partial [Trichoderma atroviride IMI 206040]
 XP_002842832.1 conserved hypothetical protein [Arthroderma otae CBS 113480] >gb|EEQ35844.1| conserved hypothetical protein [Arthroderma otae CBS 113480]
 EHK16770.1 hypothetical protein TRIVIDRAFT_135791, partial [Trichoderma virens Gv29-8]

XP_003708917.1 hypothetical protein MGG_13719 [Magnaporthe oryzae 70-15] >gb|EHA56305.1| hypothetical protein MGG_13719 [Magnaporthe oryzae 70-15]
 EME87785.1 hypothetical protein MYCFIDRAFT_129503, partial [Pseudocercospora fijiensis CIRAD86]
 EKV15847.1 Extracellular protein, putative [Penicillium digitatum Pd1]
 EKV17912.1 Extracellular protein, putative [Penicillium digitatum PHI26]
 XP_003236308.1 endoglucanase [Trichophyton rubrum CBS 118892] >gb|EGD87103.1| endoglucanase [Trichophyton rubrum CBS 118892]
 EGX93899.1 endoglucanase, putative [Cordyceps militaris CM01]
 XP_382598.1 hypothetical protein FG02422.1 [Fusarium graminearum PH-1]
 EEQ90950.1 extracellular protein [Ajellomyces dermatitidis ER-3]
 XP_002623110.1 extracellular protein [Ajellomyces dermatitidis SLH14081] >gb|EEQ71358.1| extracellular protein [Ajellomyces dermatitidis SLH14081]
 EQL36022.1 DNA-directed RNA polymerase [Ajellomyces dermatitidis ATCC 26199]
 XP_003016696.1 hypothetical protein ARB_04988 [Arthroderma benhamiae CBS 112371] >gb|EFE36051.1| hypothetical protein ARB_04988 [Arthroderma benhamiae CBS 112371]
 EFX97908.1 endoglucanase, putative [Metarhizium anisopliae ARSEF 23]
 EOD47354.1 putative endoglucanase protein [Neofusicoccum parvum UCRNP2]
 XP_002847024.1 conserved hypothetical protein [Arthroderma otae CBS 113480] >gb|EEQ31942.1| conserved hypothetical protein [Arthroderma otae CBS 113480]
 XP_003346716.1 hypothetical protein SMAC_04148 [Sordaria macrospora k-hell] >emb|CCC10919.1| unnamed protein product [Sordaria macrospora k-hell]
 XP_003018368.1 hypothetical protein TRV_07623 [Trichophyton verrucosum HKI 0517] >gb|EFE37723.1| hypothetical protein TRV_07623 [Trichophyton verrucosum HKI 0517]
 EME49904.1 hypothetical protein DOTSEDRAFT_121984, partial [Dothistroma septosporum NZE10]
 XP_003026574.1 hypothetical protein SCHCODRAFT_62159 [Schizophyllum commune H4-8] >gb|EF191671.1| hypothetical protein SCHCODRAFT_62159 [Schizophyllum commune H4-8]
 EFX201840.1 endoglucanase, putative [Metarhizium anisopliae ARSEF 23]
 XP_003025876.1 hypothetical protein SCHCODRAFT_83710 [Schizophyllum commune H4-8] >gb|EF190973.1| hypothetical protein SCHCODRAFT_83710 [Schizophyllum commune H4-8]
 EFX85839.1 endoglucanase [Metarhizium acridum CQMa 102]
 XP_003026131.1 hypothetical protein SCHCODRAFT_238682 [Schizophyllum commune H4-8] >gb|EF191228.1| hypothetical protein SCHCODRAFT_238682 [Schizophyllum commune H4-8]
 EGE82167.1 extracellular protein [Ajellomyces dermatitidis ATCC 18188]
 EEH06906.1 conserved hypothetical protein [Ajellomyces capsulatus G186AR]
 EOD53021.1 putative chitin-binding domain 3 protein [Neofusicoccum parvum UCRNP2]
 XP_001212055.1 conserved hypothetical protein [Aspergillus terreus NIH2624] >gb|EAU36151.1| conserved hypothetical protein [Aspergillus terreus NIH2624]
 EHY59273.1 DNA-directed RNA polymerase [Exophiala dermatitidis NIH/UT8656]
 XP_003029948.1 hypothetical protein SCHCODRAFT_235734 [Schizophyllum commune H4-8] >gb|EF195045.1| hypothetical protein SCHCODRAFT_235734 [Schizophyllum commune H4-8]
 XP_002340103.1 conserved hypothetical protein [Talaromyces stipitatus ATCC 10500] >gb|EED22716.1| conserved hypothetical protein [Talaromyces stipitatus ATCC 10500]
 EJP67667.1 extracellular protein [Beauveria bassiana ARSEF 2860]
 XP_002543707.1 predicted protein [Uncinocarpus reesii 1704] >gb|EEP78378.1| predicted protein [Uncinocarpus reesii 1704]
 EJP67690.1 extracellular protein [Beauveria bassiana ARSEF 2860]
 EEH50753.1 conserved hypothetical protein [Paracoccidioides brasiliensis Pb18]
 EME87736.1 hypothetical protein MYCFIDRAFT_184656 [Pseudocercospora fijiensis CIRAD86]
 EKG21662.1 endoglucanase, putative, partial [Macrophomina phaseolina MS6]
 XP_002569065.1 Pc21g20790 [Penicillium chrysogenum Wisconsin 54-1255] >emb|CAP96976.1| Pc21g20790 [Penicillium chrysogenum Wisconsin 54-1255]
 EGY22619.1 hypothetical protein VDAG_04057 [Verticillium dahliae VdLs.17]
 XP_001227363.1 hypothetical protein CHGG_09436 [Chaetomium globosum CBS 148.51] >gb|EAQ85422.1| hypothetical protein CHGG_09436 [Chaetomium globosum CBS 148.51]
 EGX88620.1 extracellular protein [Cordyceps militaris CM01]
 EEH15819.1 conserved hypothetical protein [Paracoccidioides brasiliensis Pb03]
 XP_001225174.1 hypothetical protein CHGG_07518 [Chaetomium globosum CBS 148.51] >gb|EAQ86265.1| hypothetical protein CHGG_07518 [Chaetomium globosum CBS 148.51]
 XP_003713232.1 hypothetical protein MGG_15266 [Magnaporthe oryzae 70-15] >gb|EHA53425.1| hypothetical protein MGG_15266 [Magnaporthe oryzae 70-15]
 XP_001544052.1 predicted protein [Ajellomyces capsulatus NAm1] >gb|EDN03234.1| predicted protein [Ajellomyces capsulatus NAm1]
 EMF16505.1 hypothetical protein SEPMUDRAFT_23351, partial [Sphaerulina musiva SO2202]
 EJT71157.1 endoglucanase [Gaeumannomyces graminis var. tritici R3-111a-1]
 EPQ66731.1 hypothetical protein BGT96224_1299 [Blumeria graminis f. sp. tritici 96224]
 XP_001797832.1 hypothetical protein SNOG_07498 [Phaeosphaeria nodorum SN15] >gb|EAT84964.2| hypothetical protein SNOG_07498 [Phaeosphaeria nodorum SN15]
 ELQ59175.1 hypothetical protein OOW_P131scaffold01380g2 [Magnaporthe oryzae P131]
 EJT51965.1 hypothetical protein A1Q1_06771 [Trichosporon asahii var. asahii CBS 2479] >gb|EKD02576.1| hypothetical protein A1Q2_03172 [Trichosporon asahii var. asahii CBS 8904]
 EIW68339.1 hypothetical protein TREMEDRAFT_74248 [Tremella mesenterica DSM 1558]
 XP_003656955.1 hypothetical protein THITE_2029285, partial [Thielavia terrestris NRRL 8126] >gb|AEO70619.1| hypothetical protein THITE_2029285, partial [Thielavia terrestris NRRL 8126]
 EFX95565.1 endoglucanase [Metarhizium anisopliae ARSEF 23]
 XP_002837977.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ82168.1| unnamed protein product [Tuber melanosporum]

EER38578.1 extracellular protein [Ajellomyces capsulatus H143] >gb|EGC47726.1| extracellular protein [Ajellomyces capsulatus H88]
 EON98853.1 hypothetical protein UCRPA7_5633 [Togninia minima UCRPA7]
 XP_003029539.1 hypothetical protein SCHCODRAFT_43113 [Schizophyllum commune H4-8] >gb|EF194636.1| hypothetical protein SCHCODRAFT_43113, partial [Schizophyllum commune H4-8]
 EY91408.1 endoglucanase, putative [Metarhizium acridum CQMa 102]
 CCU74383.1 endoglucanase [Blumeria graminis f. sp. hordei DH14]
 EHK21327.1 hypothetical protein TRIVIDRAFT_216235 [Trichoderma virens Gv29-8]
 XP_003850477.1 hypothetical protein MYCGRDRAFT_46444, partial [Zymoseptoria tritici IPO323] >gb|EGP85453.1| hypothetical protein MYCGRDRAFT_46444 [Zymoseptoria tritici IPO323]
 EQB52923.1 hypothetical protein CGLO_07413 [Colletotrichum gloeosporioides Cg-14]
 ELA24851.1 extracellular protein [Colletotrichum gloeosporioides Nara gc5]
 EHK19854.1 hypothetical protein TRIVIDRAFT_155830 [Trichoderma virens Gv29-8]
 XP_003195343.1 hypothetical protein CGB_G4290C [Cryptococcus gattii WM276] >gb|ADV23556.1| Conserved hypothetical protein [Cryptococcus gattii WM276]
 XP_002384586.1 conserved hypothetical protein [Aspergillus flavus NRRL3357] >gb|EED45650.1| conserved hypothetical protein [Aspergillus flavus NRRL3357]
 XP_002839220.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ83411.1| unnamed protein product [Tuber melanosporum]
 ENH80220.1 extracellular protein [Colletotrichum orbiculare MAFF 240422]
 EFQ32945.1 extracellular protein [Colletotrichum graminicola M1.001]
 XP_003352667.1 hypothetical protein SMAC_01500 [Sordaria macrospora k-hell] >emb|CCC07937.1| unnamed protein product [Sordaria macrospora k-hell]
 GAA94037.1 hypothetical protein ESQ_00684 [Mixia osmundae IAM 14324]
 EHK97268.1 putative Killer toxin subunits alpha/beta [Glarea lozoyensis 74030] >gb|EPE25430.1| Plant lectins/antimicrobial peptide [Glarea lozoyensis ATCC 20868]
 XP_002840711.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ84902.1| unnamed protein product [Tuber melanosporum]
 ELR07584.1 hypothetical protein GMDG_02632 [Pseudogymnoascus destructans 20631-21]
 XP_001225335.1 hypothetical protein CHGG_07679 [Chaetomium globosum CBS 148.51] >gb|EAQ86426.1| hypothetical protein CHGG_07679 [Chaetomium globosum CBS 148.51]
 XP_001906381.1 hypothetical protein [Podospora anserina S mat+] >emb|CAP67048.1| unnamed protein product [Podospora anserina S mat+]
 XP_003649083.1 hypothetical protein THITE_120654 [Thielavia terrestris NRRL 8126] >gb|AEO62747.1| hypothetical protein THITE_120654 [Thielavia terrestris NRRL 8126]
 XP_003029201.1 hypothetical protein SCHCODRAFT_16695 [Schizophyllum commune H4-8] >gb|EF194298.1| hypothetical protein SCHCODRAFT_16695 [Schizophyllum commune H4-8]
 XP_001827392.2 endoglucanase [Aspergillus oryzae RIB40]
 XP_002838243.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ82434.1| unnamed protein product [Tuber melanosporum]
 XP_002835201.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ79322.1| unnamed protein product [Tuber melanosporum]
 XP_003836919.1 predicted protein [Leptosphaeria maculans JN3] >emb|CBX93554.1| predicted protein [Leptosphaeria maculans JN3]
 XP_001911580.1 hypothetical protein [Podospora anserina S mat+] >emb|CAP73406.1| unnamed protein product [Podospora anserina S mat+]
 EMC94923.1 hypothetical protein BAUCODRAFT_518344 [Baudoinia compniacensis UAMH 10762]
 XP_001594978.1 hypothetical protein SS1G_04786 [Sclerotinia sclerotiorum 1980] >gb|EDO02310.1| hypothetical protein SS1G_04786 [Sclerotinia sclerotiorum 1980 UF-70]
 XP_571909.1 hypothetical protein [Cryptococcus neoformans var. neoformans JEC21] >gb|AAW44602.1| conserved hypothetical protein [Cryptococcus neoformans var. neoformans JEC21]
 XP_774319.1 hypothetical protein CNBG3000 [Cryptococcus neoformans var. neoformans B-3501A] >gb|EAL19672.1| hypothetical protein CNBG3000 [Cryptococcus neoformans var. neoformans B-3501A]
 AFR96628.1 hypothetical protein CNAG_03405 [Cryptococcus neoformans var. grubii H99]
 EQB58862.1 hypothetical protein CGLO_00836 [Colletotrichum gloeosporioides Cg-14]
 EKG09287.1 hypothetical protein MPH_13695 [Macrophomina phaseolina MS6]
 XP_003717516.1 hypothetical protein MGG_13619 [Magnaporthe oryzae 70-15] >gb|EHA51197.1| hypothetical protein MGG_13619 [Magnaporthe oryzae 70-15]
 EKD12205.1 RNA polymerase Rpb1 repeat domain protein [Marssonina brunnea f. sp. 'multigermtubi' MB_m1]
 EGX47272.1 hypothetical protein AOL_s00091g16 [Arthrotrichy oligospora ATCC 24927]
 XP_003667236.1 hypothetical protein MYCTH_2312852 [Myceliophthora thermophila ATCC 42464] >gb|AEO61991.1| hypothetical protein MYCTH_2312852 [Myceliophthora thermophila ATCC 42464]
 XP_003649387.1 hypothetical protein THITE_2038036 [Thielavia terrestris NRRL 8126] >gb|AEO63051.1| hypothetical protein THITE_2038036 [Thielavia terrestris NRRL 8126]
 EJT76146.1 hypothetical protein GGTG_06068 [Gaeumannomyces graminis var. tritici R3-111a-1]
 XP_003050224.1 hypothetical protein NECHADRAFT_85135 [Nectria haematococca mpVI 77-13-4] >gb|EEU44511.1| hypothetical protein NECHADRAFT_85135 [Nectria haematococca mpVI 77-13-4]
 ELQ61523.1 hypothetical protein OOW_P131scaffold01178g3 [Magnaporthe oryzae P131]
 ELA32984.1 C6 transcription factor [Colletotrichum gloeosporioides Nara gc5]
 ELR08410.1 hypothetical protein GMDG_03199 [Pseudogymnoascus destructans 20631-21]
 XP_001549934.1 hypothetical protein BC1G_11826 [Botryotinia fuckeliana B05.10] >gb|EMR80832.1| putative carbohydrate-binding module family 18 protein [Botryotinia fuckeliana BcDW1]
 XP_001797251.1 hypothetical protein SNOG_06890 [Phaeosphaeria nodorum SN15] >gb|EAT85541.2| hypothetical protein SNOG_06890 [Phaeosphaeria nodorum SN15]
 CCD54857.1 carbohydrate-Binding Module family 18 protein [Botryotinia fuckeliana T4]
 EGO52047.1 hypothetical protein NEUTE1DRAFT_53322 [Neurospora tetrasperma FGSC 2508]
 XP_964365.1 hypothetical protein NCU00822 [Neurospora crassa OR74A] >gb|EAA35129.1| predicted protein [Neurospora crassa OR74A]
 EPS44930.1 hypothetical protein H072_1087 [Dactylellina haptotyla CBS 200.50]
 XP_003709778.1 endoglucanase [Magnaporthe oryzae 70-15] >gb|EHA57166.1| endoglucanase [Magnaporthe oryzae 70-15]

ELQ35674.1 hypothetical protein OOU_Y34scaffold00694g10 [Magnaporthe oryzae Y34]
 EGZ76857.1 hypothetical protein NEUTE2DRAFT_134115 [Neurospora tetrasperma FGSC 2509]
 ELQ36644.1 hypothetical protein OOU_Y34scaffold00649g27 [Magnaporthe oryzae Y34]
 XP_001242938.1 hypothetical protein CIMG_06834 [Coccidioides immitis RS]
 EIT75494.1 hypothetical protein Ao3042_08433 [Aspergillus oryzae 3.042]
 BAE66259.1 unnamed protein product [Aspergillus oryzae RIB40]
 EGX43927.1 hypothetical protein AOL_s00210g374 [Arthrotrichum oligospora ATCC 24927]
 EMR68381.1 putative extracellular protein [Eutypa lata UCREL1]
 XP_001220488.1 hypothetical protein CHGG_01267 [Chaetomium globosum CBS 148.51] >gb|EAQ93032.1| hypothetical protein CHGG_01267 [Chaetomium globosum CBS 148.51]
 EMR61516.1 putative extracellular protein [Eutypa lata UCREL1]
 XP_002838570.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ82761.1| unnamed protein product [Tuber melanosporum]
 EON64769.1 hypothetical protein W97_04002 [Coniosporium apollinis CBS 100218]
 EKD13625.1 endoglucanase [Marssonina brunnea f. sp. 'multigermtubi' MB_m1]
 XP_003712361.1 hypothetical protein MGG_04880 [Magnaporthe oryzae 70-15] >gb|EHA52554.1| hypothetical protein MGG_04880 [Magnaporthe oryzae 70-15]
 EMS24852.1 endoglucanase [Rhodosporidium toruloides NP11]
 EQB56592.1 hypothetical protein CGLO_03386 [Colletotrichum gloeosporioides Cg-14]
 ELQ68311.1 hypothetical protein OOW_P131scaffold00255g13 [Magnaporthe oryzae P131]
 GAA95388.1 hypothetical protein ESQ_02042 [Mixia osmundae IAM 14324]
 XP_002841287.1 hypothetical protein [Tuber melanosporum Mel28] >emb|CAZ85478.1| unnamed protein product [Tuber melanosporum]
 EJT74193.1 hypothetical protein GGTG_08038 [Gaeumannomyces graminis var. tritici R3-111a-1]
 ELQ39898.1 hypothetical protein OOU_Y34scaffold00468g1, partial [Magnaporthe oryzae Y34]
 XP_003011652.1 hypothetical protein ARB_02206 [Arthroderma benhamiae CBS 112371] >gb|EFE44783.1| hypothetical protein TRV_00455 [Trichophyton verrucosum HKI 0517]
 XP_003231393.1 hypothetical protein TERG_08178 [Trichophyton rubrum CBS 118892] >gb|EGD91962.1| hypothetical protein TERG_08178 [Trichophyton rubrum CBS 118892]
 XP_002842573.1 conserved hypothetical protein [Arthroderma otae CBS 113480] >gb|EEQ28554.1| conserved hypothetical protein [Arthroderma otae CBS 113480]
 EGE06977.1 endoglucanase [Trichophyton equinum CBS 127.97]
 XP_003171247.1 hypothetical protein MGYG_07246 [Arthroderma gypseum CBS 118893] >gb|EFR04239.1| hypothetical protein MGYG_07246 [Arthroderma gypseum CBS 118893]
 EGD99938.1 hypothetical protein TEGS_07268 [Trichophyton tonsurans CBS 112818]
 XP_003658966.1 hypothetical protein MYCTH_2295431 [Myceliophthora thermophila ATCC 42464] >gb|AEO53721.1| hypothetical protein MYCTH_2295431 [Myceliophthora thermophila ATCC 42464]
 ADK37848.1 putative endoglucanase precursor [Entomophthora planchoniana]
 XP_00267479.1 predicted protein [Naegleria gruberi] >gb|EFC41435.1| predicted protein [Naegleria gruberi]
 XP_002556576.1 Pc03g00070 [Penicillium chrysogenum Wisconsin 54-1255] >emb|CAP78969.1| Pc03g00070 [Penicillium chrysogenum Wisconsin 54-1255]
 CCF40911.1 hypothetical protein CH063_02459 [Colletotrichum higginsianum]
 EON65846.1 hypothetical protein W97_05088 [Coniosporium apollinis CBS 100218]
 XP_003854534.1 hypothetical protein MYCGRDRAFT_38707 [Zymoseptoria tritici IPO323] >gb|EGP89510.1| hypothetical protein MYCGRDRAFT_38707 [Zymoseptoria tritici IPO323]
 ENH82340.1 glycoside hydrolase family 92 [Colletotrichum orbiculare MAFF 240422]
 XP_001541987.1 predicted protein [Ajellomyces capsulatus NAM1] >gb|EDN05555.1| predicted protein [Ajellomyces capsulatus NAM1]
 EEH09607.1 conserved hypothetical protein [Ajellomyces capsulatus G186AR]
 EGC44079.1 endoglucanase [Ajellomyces capsulatus H88]
 EER37945.1 endoglucanase [Ajellomyces capsulatus H143]
 EEQ89606.1 endoglucanase [Ajellomyces dermatitidis ER-3]
 ELQ69881.1 hypothetical protein OOW_P131scaffold00107g1 [Magnaporthe oryzae P131]
 EMC92258.1 hypothetical protein BAUCODRAFT_283706 [Baudoinia compniacensis UAMH 10762]
 EGE85732.1 endoglucanase [Ajellomyces dermatitidis ATCC 18188] >gb|EQL34786.1| hypothetical protein BDFG_03482 [Ajellomyces dermatitidis ATCC 26199]
 XP_002622207.1 endoglucanase [Ajellomyces dermatitidis SLH14081] >gb|EEQ72616.1| endoglucanase [Ajellomyces dermatitidis SLH14081]
 GAA98995.1 hypothetical protein ESQ_05684 [Mixia osmundae IAM 14324]
 EOD49695.1 putative chitin-binding domain 3 protein [Neofusicoccum parvum UCRNP2]
 EMD66417.1 hypothetical protein COCSADRAFT_24530 [Bipolaris sorokiniana ND90Pr]
 EMD84831.1 hypothetical protein COCHEDRAFT_1051068, partial [Bipolaris maydis C5] >gb|EMD96369.1| hypothetical protein COCHEDRAFT_1046206, partial [Bipolaris maydis C5]
 XP_003843789.1 predicted protein [Leptosphaeria maculans JN3] >emb|CBY00310.1| predicted protein [Leptosphaeria maculans JN3]
 EME84178.1 hypothetical protein MYCFIDRAFT_195305 [Pseudocercospora fijiensis CIRAD86]
 EMD65813.1 hypothetical protein COCSADRAFT_48501, partial [Bipolaris sorokiniana ND90Pr]
 EEH16034.1 conserved hypothetical protein [Paracoccidioides brasiliensis Pb03]
 EMS21926.1 endoglucanase [Rhodosporidium toruloides NP11]

EMD85633.1 hypothetical protein COCHEDRAFT_1038311, partial [Bipolaris maydis C5] >gb|ENH99065.1| hypothetical protein COCC4DRAFT_95266, partial [Bipolaris maydis ATCC 48331]
EEH50505.1 conserved hypothetical protein [Paracoccidioides brasiliensis Pb18]
EOA90842.1 hypothetical protein SETTUDRAFT_36346 [Setosphaeria turcica Et28A]
EAS32093.2 hypothetical protein CIMG_03033 [Coccidioides immitis RS]
EON62244.1 hypothetical protein W97_01465 [Coniosporium apollinis CBS 100218]
EFW18701.1 conserved hypothetical protein [Coccidioides posadasii str. Silveira]
XP_001243592.1 hypothetical protein CIMG_03033 [Coccidioides immitis RS]
XP_003069064.1 hypothetical protein CPC735_010980 [Coccidioides posadasii C735 delta SOWgp]
EMD86612.1 hypothetical protein COCHEDRAFT_22861 [Bipolaris maydis C5] >gb|ENI03016.1| hypothetical protein COCC4DRAFT_62842 [Bipolaris maydis ATCC 48331]
XP_003843057.1 hypothetical protein LEMA_P088170.1 [Leptosphaeria maculans JN3] >emb|CBX99578.1| hypothetical protein LEMA_P088170.1 [Leptosphaeria maculans JN3]
EOA83923.1 hypothetical protein SETTUDRAFT_61484, partial [Setosphaeria turcica Et28A]
XP_003297367.1 hypothetical protein PTT_07742 [Pyrenophora teres f. teres 0-1] >gb|EFQ94536.1| hypothetical protein PTT_07742 [Pyrenophora teres f. teres 0-1]
EOD47246.1 putative endoglucanase protein [Neofusicoccum parvum UCRNP2]
XP_001940795.1 conserved hypothetical protein [Pyrenophora tritici-repentis Pt-1C-BFP] >gb|EDU43514.1| conserved hypothetical protein [Pyrenophora tritici-repentis Pt-1C-BFP]
EMD65810.1 hypothetical protein COCSADRAFT_355161 [Bipolaris sorokiniana ND90Pr]
XP_001792664.1 hypothetical protein SNOG_02045 [Phaeosphaeria nodorum SN15] >gb|EAT90257.1| hypothetical protein SNOG_02045 [Phaeosphaeria nodorum SN15]
EKG12228.1 hypothetical protein MPH_10635 [Macrophomina phaseolina MS6]
EKG09975.1 hypothetical protein MPH_12937 [Macrophomina phaseolina MS6]
EME45691.1 hypothetical protein DOTSEDRAFT_79599 [Dothistroma septosporum NZE10]
EMF10416.1 hypothetical protein SEPMUDRAFT_143102 [Sphaerulina musiva SO2202]
XP_003305986.1 hypothetical protein PTT_18989 [Pyrenophora teres f. teres 0-1] >gb|EFQ85911.1| hypothetical protein PTT_18989 [Pyrenophora teres f. teres 0-1]
XP_002789291.1 conserved hypothetical protein [Paracoccidioides sp. 'lutzii' Pb01] >gb|EEH39527.1| conserved hypothetical protein [Paracoccidioides sp. 'lutzii' Pb01]
EOA90850.1 hypothetical protein SETTUDRAFT_101501 [Setosphaeria turcica Et28A]
XP_001261280.1 hypothetical protein NFIA_093510 [Neosartorya fischeri NRRL 181] >gb|EAW19383.1| hypothetical protein NFIA_093510 [Neosartorya fischeri NRRL 181]
ENH99059.1 hypothetical protein COCC4DRAFT_66801 [Bipolaris maydis ATCC 48331]
EMD84834.1 hypothetical protein COCHEDRAFT_1121170 [Bipolaris maydis C5] >gb|EMD96372.1| hypothetical protein COCHEDRAFT_1152415 [Bipolaris maydis C5]
EOA85932.1 hypothetical protein SETTUDRAFT_110770 [Setosphaeria turcica Et28A]
EOA83921.1 hypothetical protein SETTUDRAFT_92647 [Setosphaeria turcica Et28A]
EMD66420.1 hypothetical protein COCSADRAFT_84949 [Bipolaris sorokiniana ND90Pr]
EOA85398.1 hypothetical protein SETTUDRAFT_164047 [Setosphaeria turcica Et28A]
EMD94762.1 hypothetical protein COCHEDRAFT_1167926 [Bipolaris maydis C5] >gb|ENI01527.1| hypothetical protein COCC4DRAFT_176644 [Bipolaris maydis ATCC 48331]
EMD58702.1 hypothetical protein COCSADRAFT_41806 [Bipolaris sorokiniana ND90Pr]
EMD85627.1 hypothetical protein COCHEDRAFT_75573 [Bipolaris maydis C5]